

09/834,271

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(FILE 'HOME' ENTERED AT 14:23:15 ON 23 FEB 2006)

FILE 'MEDLINE, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH, HCAPLUS, NTIS, LIFESCI' ENTERED AT 14:23:47 ON 23 FEB 2006

L1 395086 S BACILLUS  
L2 560 S CRYIIIA  
L3 516 S L1 AND L2  
L4 635 S "TTGACA" OR "TATAAT"  
L5 13 S L3 AND L4  
L6 6 DUP REM L5 (7 DUPLICATES REMOVED)  
L7 7 S L2 AND "CONSENSUS PROMOTER"  
L8 5 DUP REM L7 (2 DUPLICATES REMOVED)  
E WIDNER W/AU  
L9 119 S E3-E9  
E SLOMA A/AU  
L10 124 S E3  
E THOMAS M D/AU  
L11 404 S E3  
L12 632 S L9 OR L10 OR L11  
L13 8 S L2 AND L12  
L14 6 DUP REM L13 (2 DUPLICATES REMOVED)

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IPC reform  
NEWS 8 DEC 23 New IPC8 SEARCH, DISPLAY, and SELECT fields in USPATFULL/  
USPAT2  
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NEWS 14 JAN 31 Monthly current-awareness alert (SDI) frequency  
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visualization results  
NEWS 16 FEB 22 Status of current WO (PCT) information on STN  
NEWS 17 FEB 22 The IPC thesaurus added to additional patent databases on STN  
NEWS 18 FEB 22 Updates in EPFULL; IPC 8 enhancements added  
  
NEWS EXPRESS FEBRUARY 15 CURRENT VERSION FOR WINDOWS IS V8.01a,  
CURRENT MACINTOSH VERSION IS V6.0c(ENG) AND V6.0Jc(JP),  
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=> s bacillus  
L1 395086 BACILLUS

=> s cryIIIA  
L2 560 CRYIIIA

=> s l1 and l2  
L3 516 L1 AND L2

=> s "TTGACA" or "TATAAT"  
L4 635 "TTGACA" OR "TATAAT"

=> s l3 and l4  
L5 13 L3 AND L4

=> dup rem l5  
PROCESSING COMPLETED FOR L5  
L6 6 DUP REM L5 (7 DUPLICATES REMOVED)

=> d 1-6 ibib ab

L6 ANSWER 1 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN  
DUPLICATE 1

ACCESSION NUMBER: 2005-30339 BIOTECHDS

TITLE: Producing a hyaluronic acid, by cultivating *Bacillus*  
cell comprising a nucleic acid construct comprising a variant  
amyL promoter, a consensus promoter, and a cryIIIA  
promoter, and isolating hyaluronic acid from the cultivation  
medium;  
production of recombinant hyaluronic acid from a

**Bacillus having a triple promoter useful for a tissue engineering application**

AUTHOR: WIDNER W; SLOMA A; THOMAS M; TANG M  
PATENT ASSIGNEE: NOVOZYMES BIOPOLYMER AS  
PATENT INFO: US 2005221446 6 Oct 2005  
APPLICATION INFO: US 2005-96190 31 Mar 2005  
PRIORITY INFO: US 2005-96190 31 Mar 2005; US 2004-558507 31 Mar 2004  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
OTHER SOURCE: WPI: 2005-722702 [74]

**AB DERWENT ABSTRACT:**

**NOVELTY** - Producing a hyaluronic acid comprises cultivating a **Bacillus** cell in a medium for the production of the hyaluronic acid, where the **Bacillus** cell comprises a nucleic acid construct comprising a triple promoter comprising a variant amyL promoter, a consensus promoter, and a **cryIIIA** promoter, and isolating the hyaluronic acid from the cultivation medium.

**DETAILED DESCRIPTION** - Producing a hyaluronic acid comprises: (A) cultivating a **Bacillus** cell in a medium for the production of the hyaluronic acid, where the **Bacillus** cell comprises a nucleic acid construct comprising a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of a fully defined 614 bp sequence (SEQ ID NO. 1), a consensus promoter having the sequence **TTGACA** for the "-35" region and **TATAAT** for the "-10" region, and a **cryIIIA** promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of the hyaluronic acid; and (B) isolating the hyaluronic acid from the cultivation medium. **INDEPENDENT CLAIMS** are also included for: (1) a **Bacillus** cell comprising a nucleic acid construct which comprises: (a) a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of SEQ ID NO. 1, a consensus promoter having the sequence **TTGACA** for the "-35" region and **TATAAT** for the "-10" region, and a **cryIIIA** promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of a hyaluronic acid, and optionally (b) an mRNA processing/stabilizing sequence located downstream of the triple promoter and upstream of the one or more coding sequences involved in the biosynthesis of the hyaluronic acid; (2) producing a selectable marker-free mutant of a **Bacillus** cell; and (3) a selectable marker-free mutant of a **Bacillus** cell obtained by the method above.

**WIDER DISCLOSURE** - Also disclosed are: (1) methods for obtaining a **Bacillus** host cell; and (2) nucleic acid constructs comprising a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of SEQ ID NO. 1, a consensus promoter having the sequence **TTGACA** for the "-35" region and **TATAAT** for the "-10" region, and a **cryIIIA** promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of a hyaluronic acid.

**BIOTECHNOLOGY** - Preferred Method: In producing a hyaluronic acid, the variant amyL promoter is SEQ ID NO. 1. The consensus promoter is obtained from the **Bacillus** amyloliquefaciens alpha-amylase gene (amyQ), where the consensus amyQ promoter has the nucleotide sequence comprising fully defined 185 bp sequence (SEQ ID NO. 42 or 43). The nucleic acid construct further comprises an mRNA processing/stabilizing sequence located downstream of the triple promoter and upstream of the one or more coding sequences involved in the biosynthesis of the hyaluronic acid. Preferably, the one or more coding sequences involved in the biosynthesis of the hyaluronic acid are selected from a hyaluronan synthase, UDP-glucose 6-dehydrogenase, UDP-glucose pyrophosphorylase, UDP-N-acetylglucosamine pyrophosphorylase, glucose-6-phosphate isomerase, hexokinase, phosphoglucomutase, amidotransferase, mutase, or acetyl

transferase gene. Producing a selectable marker-free mutant of a *Bacillus* cell comprises deleting a selectable marker gene of the *Bacillus* cell. The *Bacillus* cell contains no foreign selectable marker gene.

USE - The method is useful for producing a hyaluronic acid. Hyaluronic acid is useful in eye and joint surgery. Products of hyaluronic acid are also useful in orthopedics, rheumatology, and in dermatology.

EXAMPLE - No relevant example given. (94 pages)

L6 ANSWER 2 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN  
DUPLICATE 2

ACCESSION NUMBER: 2004-02033 BIOTECHDS

TITLE: Generating an expression library of polynucleotides by introducing the linear integration cassette into the host cell and selecting or screening for host cells that produce the polypeptides of interest;  
protein library screening using homologous recombination

AUTHOR: BJORNVAD M E; JORGENSEN P L; HANSEN P K

PATENT ASSIGNEE: NOVOZYMES AS

PATENT INFO: WO 2003095658 20 Nov 2003

APPLICATION INFO: WO 2003-DK301 7 May 2003

PRIORITY INFO: DK 2002-682 7 May 2002; DK 2002-682 7 May 2002

DOCUMENT TYPE: Patent

LANGUAGE: English

AB DERWENT ABSTRACT:

NOVELTY - Generating an expression library of polynucleotides integrated by homologous recombination into the genome of a competent Gram-positive bacterium host cell comprises: (a) providing a non-replicating linear integration cassette; and (b) introducing the linear integration cassette into the host cell and selecting or screening for host cells that produce the polypeptides of interest.

DETAILED DESCRIPTION - Generating an expression library of polynucleotides integrated by homologous recombination into the genome of a competent Gram-positive bacterium host cell comprises: (a) providing a non-replicating linear integration cassette; and (b) introducing the linear integration cassette into the host cell and selecting or screening for host cells that produce the polypeptides of interest. The cassette comprises: (1) a polynucleotide encoding one or more polypeptides of interest; (2) a 5' flanking polynucleotide segment upstream of the polynucleotide of (1) and comprising a first homologous region located in the 3' end of the segment; and (3) a 3' flanking polynucleotide segment downstream of the polynucleotide of (1) and comprising a second homologous region located in the 5' end of the segment. The first and second homologous regions are at least 500, 1000 or 1500 bp, each of which has a sequence identity of at least 80, 85, 90 or 95-100% with a region of the host cell genom. INDEPENDENT CLAIMS are also included for the following: (1) a non-replicating linear Gram-positive host cell integration cassette; and (2) a method of producing a polypeptide of interest.

BIOTECHNOLOGY - Preferred Method: Generating an expression library of polynucleotides integrated by homologous recombination into the genome of a competent Gram-positive bacterium host cell comprises the additional step carried out between steps (1) and (2) that comprises introducing the plasmid into an intermediate *Escherichia coli* host cell and propagating it by replication. The integration cassette comprises: (1) an mRNA processing/stabilizing sequence derived from *cryIIIA*-gene; (2) a terminator downstream of the polynucleotide encoding the polypeptides of interest; and (3) a multiple cloning site with at least one recognition site for a restriction nuclease. It further comprises a marker gene located between the 5' and 3' flanking segments and at least one promoter that is a consensus promoter having the sequence TTGACA for the -35 region and TATAAT for the -10 region, and/or which is derived from *amyL*, *amyQ*, *amyM*, *cryIIIA*,

dagA, aprH, penP, sacB, spol, tac, xylA or xylB. The promoter is located between the flanking segments and is operably linked to the polynucleotide encoding one or more polypeptides of interest. Each of the 5' and 3' flanking polynucleotide segments comprises at least 500, 1000, 1500 or 2000 bp of non-homologous polynucleotides located in the 5' and 3' end of the 5' and 3' flanking segments, respectively. The promoter is one that results in that the host cells produce the polypeptides of interest in a yield of at least 10 mg/L. The polynucleotide comprises natural, synthetic or a library of shuffled or recombined homologs or variants of a gene or operon, provided by DNA breeding or DNA shuffling. The polypeptides of interest comprise enzymes, proteins or antimicrobial peptides. The enzymes are involved in the biosynthesis of hyaluronic acid. The Gram-positive host cell is *Bacillus subtilis*. The homologous region of the 5' and/or the 3' flanking segment is comprised in the yfmD-yfmC-yfmB-yfmA-pelB-yf1S-citS region of the *Bacillus subtilis* genome or in the cryIIIa promoter. The non-replicating linear integration cassette is comprised in a plasmid and introduced into the host cell. The plasmid is capable of replicating in an *Escherichia coli* host cell but not in a *Bacillus* host cell. Producing a polypeptide of interest comprises culturing Gram-positive host cells comprising the integration cassette integrated into its genome, under conditions promoting expression of the polypeptide of interest. The method further comprises isolating and/or purifying the polypeptide of interest.

USE - The method is useful in generating an expression library of polynucleotides integrated by homologous recombination into the genome of a competent Gram-positive bacterium host cell (claimed). (55 pages)

L6 ANSWER 3 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-08483 BIOTECHDS

TITLE: Production of a secreted polypeptide having L-asparaginase activity for treating leukemia, comprises cultivating a host cell comprising a nucleic acid having a sequence encoding a secretory signal peptide linked to a second sequence; vector-mediated enzyme gene transfer and expression in host cell for recombinant protein production, amino acid preparation and disease therapy

AUTHOR: THOMAS M D; SLOMA A

PATENT ASSIGNEE: NOVOZYMES BIOTECH INC

PATENT INFO: US 2003186380 2 Oct 2003

APPLICATION INFO: US 2003-406025 1 Apr 2003

PRIORITY INFO: US 2003-406025 1 Apr 2003; US 2002-369192 1 Apr 2002

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: WPI: 2004-088916 [09]

AB DERWENT ABSTRACT:

NOVELTY - Producing a secreted polypeptide having L-asparaginase activity, comprises cultivating a host cell containing a nucleic acid construct having a sequence encoding a secretory signal peptide linked to a second sequence encoding the polypeptide having L-asparaginase activity, where the signal peptide directs the polypeptide into the cell's secretory pathway, and recovering the secreted polypeptide.

DETAILED DESCRIPTION - An INDEPENDENT CLAIM is also included for a recombinant expression vector comprising the nucleic acid construct.

BIOTECHNOLOGY - Preferred Material: The polypeptide is encoded by a nucleic acid sequence contained in plasmid pCR2.1-yccc, which is contained in *Escherichia coli* (NRRL B-30558). The consensus promoter is obtained from a promoter obtained from the *E. coli* lac operon *Streptomyces coelicolor* agarase gene (dagA), *Bacillus clausii* alkaline protease gene (aprH), *B. licheniformis* alkaline protease gene (subtilisin Carlsberg gene), *B. subtilis* levansucrase gene (sacB), *B. subtilis* alpha-amylase gene (amyE), *B. licheniformis* alpha-amylase gene (amyL), *B. stearothermophilus* maltogenic amylase gene (amyM), *B. licheniformis* penicillinase gene (penP), *B. subtilis* xylA and xylB genes,

*B. thuringiensis* subsp. *tenebrionis* **CryIIIA** gene ( **cryIIIA**) or its portions, or preferably *B. amyloliquefaciens* alpha-amylase gene (**amyQ**). The mRNA processing/stabilizing sequence is the **cryIIIA** mRNA processing/stabilizing sequence. The **bacillus** cell is *B. alkalophilus*, *B. amyloliquefaciens*, *B. brevis*, *B. circulans*, *B. clausii*, *B. coagulans*, *B. lautus*, *B. lentus*, *B. licheniformis*, *B. megaterium*, *B. stearothermophilus*, *B. subtilis*, or *B. thuringiensis*. Preferred Component: The nucleic acid construct comprises a tandem promoter, in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding a polypeptide, and optionally an mRNA processing stabilizing sequence located downstream of the tandem promoter and upstream of the second nucleic acid sequence, encoding the polypeptide having L-asparaginase activity. It comprises a consensus promoter having the sequence **TTGACA** for the 35 region, and **TATAAT** for the 10 region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and an mRNA processing stabilizing sequence located downstream of the consensus promoter, and upstream of the second nucleic acid sequence encoding the polypeptide having L-asparagine activity. The consensus promoter is obtained from any bacterial or a **bacillus** promoter. The nucleic acid construct comprises a ribosome binding site sequence heterologous to the host cell.

ACTIVITY - Cytostatic.

MECHANISM OF ACTION - L-asparaginase.

USE - The invention is for production of secreted polypeptide having L-asparaginase activity, for use in producing L-aspartate from L-asparagine. The inventive secreted polypeptide is useful for treatment of leukemia, e.g. acute lymphocytic leukemia.

ADVANTAGE - The invention achieves secretion of L-asparaginase enabling easy recovery and purification, high expression constructs for producing the L-asparaginase in high amounts, and the use of host cells for production that have generally regarded as safe status.

EXAMPLE - *B. subtilis* strains MDT51 and MDT52 were grown in Lactobacilli MRS Broth (RTM; 50 ml) at 37degreesC, and 250 revolutions/minute (rpm) for 24 hours. Supernatants were recovered by centrifugation at 7000 rpm for 5 minutes. A prominent band corresponding to a protein of the expected size for mature L-asparagine (37 kDa; amino acids 24-375) was observed in the MDT51 sample, but not in the MDT52 sample. (22 pages)

L6 ANSWER 4 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-04169 BIOTECHDS

TITLE: Producing a polypeptide comprises cultivating a **Bacillus** cell in a medium conducive to the production of the polypeptide, where the **Bacillus** cell comprises a nucleic acid construct comprising a tandem promoter;  
involving vector-mediated gene transfer and expression in host cell for use as a selectable marker

AUTHOR: WIDNER W; SLOMA A; THOMAS M D

PATENT ASSIGNEE: NOVOZYMES BIOTECH INC

PATENT INFO: US 2003170876 11 Sep 2003

APPLICATION INFO: US 2001-834271 12 Apr 2001

PRIORITY INFO: US 2001-834271 12 Apr 2001; US 1998-31442 26 Feb 1998

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: WPI: 2003-898275 [82]

AB DERWENT ABSTRACT:

NOVELTY - Producing a polypeptide comprises cultivating a **Bacillus** cell in a medium conducive to the production of the polypeptide, where the **Bacillus** cell comprises a nucleic acid construct comprising a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a nucleic acid sequence encoding the polypeptide, and isolating the polypeptide from the

cultivation medium.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are included for the following: (1) a *Bacillus* cell comprising a nucleic acid construct comprising a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding a polypeptide, and optionally an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide; (2) a method for obtaining a *Bacillus* host cell by introducing into a *Bacillus* cell the nucleic acid construct cited above; (3) a method for producing a selectable marker-free mutant of a *Bacillus* cell by deleting a selectable marker gene of the *Bacillus* cell; and (4) a selectable marker-free mutant of a *Bacillus* cell.

BIOTECHNOLOGY - Preferred Method: In producing a polypeptide, the nucleic acid construct further comprises an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide. The tandem promoter comprises two or more bacterial promoter sequences, which are obtained from one or more *Bacillus* genes. The tandem promoter comprises the amyQ promoter, a consensus promoter having the sequence TTGACA for the -35 region and TATAAT from the -10 region, the amyL promoter, and/or the cryIIIA promoter. The tandem promoter comprises two copies of the amyQ, amyL or cryIIIA promoter. The two or more promoter sequences of the tandem promoter simultaneously promote the transcription of the nucleic acid sequence. The one or more of the two or more promoter sequences of the tandem promoter promote the transcription of the nucleic acid sequence at different stages of growth of the *Bacillus* cell. The mRNA processing/stabilizing sequence is the cryIIIA or SP82 mRNA processing/stabilizing sequence, which generates mRNA transcripts essentially of the same size. The *Bacillus* cell contains one or more copies of the nucleic acid construct. The nucleic acid construct further comprises a selectable marker gene. The nucleic acid sequence encodes a polypeptide heterologous to the *Bacillus* cell. The polypeptide is a hormone or its variant, enzyme, receptor or its portion, antibody or its portion, or reporter. The enzyme is an oxidoreductase, transferase, hydrolase, lyase, isomerase or ligase. The enzyme is preferably an aminopeptidase, amylase, carbohydrase, carboxypeptidase, catalase, cellulase, chitinase, cutinase, cyclodextrin glycosyltransferase, deoxyribonuclease, esterase, alpha-galactosidase, beta-galactosidase, glucoamylase, alpha-glucosidase, beta-glucosidase, invertase, laccase, lipase, mannosidase, mutanase, oxidase, a pectinolytic enzyme, peroxidase, phytase, polyphenoloxidase, proteolytic enzyme, ribonuclease, transglutaminase or xylanase. The nucleic acid sequence is contained in the chromosome of the *Bacillus* cell. The *Bacillus* host cell is *Bacillus alkalophilus*, *Bacillus amyloliquefaciens*, *Bacillus brevis*, *Bacillus brevis*, *Bacillus circulans*, *Bacillus clausii*, *Bacillus coagulans*, *Bacillus firmus*, *Bacillus lautus*, *Bacillus lentus*, *Bacillus licheniformis*, *Bacillus megaterium*, *Bacillus pumilus*, *Bacillus stercorophilus*, *Bacillus subtilis*, or *Bacillus thuringiensis*. This method alternatively comprises cultivating a *Bacillus* cell in a medium conducive for the production of the polypeptide, where the *Bacillus* cell comprises a nucleic acid construct comprising a consensus promoter having the sequence TTGACA for the -35 region and TATAAT for the -10 region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and an mRNA processing/stabilizing sequence located downstream of the consensus promoter and upstream of the nucleic acid sequence encoding the polypeptide; and isolating the polypeptide from the cultivation medium. The consensus promoter is obtained from any bacterial promoter, preferably a *Bacillus*



promoter. Preferred Cell: The **Bacillus** cell comprises a nucleic acid construct that further comprises a selectable marker gene. The cell can also contain no selectable marker gene.

USE - The methods are useful for producing a polypeptide in a **Bacillus** cell, and for producing a selectable marker-free mutant of a **Bacillus** cell.

EXAMPLE - No relevant example given. (57 pages)

L6 ANSWER 5 OF 6 MEDLINE on STN DUPLICATE 3  
ACCESSION NUMBER: 2001393416 MEDLINE  
DOCUMENT NUMBER: PubMed ID: 11234961  
TITLE: Construction of protein overproducer strains in **Bacillus subtilis** by an integrative approach.  
AUTHOR: Jan J; Valle F; Bolivar F; Merino E  
CORPORATE SOURCE: Departamento de Microbiologia Molecular, Instituto de Biotecnologia, Universidad Nacional Autonoma de Mexico, Cuernavaca, Morelos.  
SOURCE: Applied microbiology and biotechnology, (2001 Jan) 55 (1) 69-75.  
Journal code: 8406612. ISSN: 0175-7598.  
PUB. COUNTRY: Germany: Germany, Federal Republic of  
DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)  
LANGUAGE: English  
FILE SEGMENT: Priority Journals  
ENTRY MONTH: 200107  
ENTRY DATE: Entered STN: 20010716  
Last Updated on STN: 20010716  
Entered Medline: 20010712  
AB We evaluated the effect of several genetic factors reported as having a role in the induction of the expression of significant levels of recombinant protein in **Bacillus subtilis**. We utilized the beta-galactosidase reporter protein from *Escherichia coli* as our model for measuring the overproduction of heterologous proteins in *B. subtilis*. The lacZ gene was expressed in *B. subtilis* using the regulatory region of the subtilisin gene aprE. In this study, we considered factors known to modulate the transcription and translation initiation rates and genetic and mRNA stability. We also consider the effects of different genetic backgrounds, such as degU32 and hpr2, that until now have been studied independently. By changing the native -35 promoter box to the consensus TTGACA sequence of the aprE promoter, a significant 100-fold increase in the beta-galactosidase activity was obtained. On the other hand, changes such as the GTG to ATG start codon, the construction of a consensus AAGGAGG ribosome binding site, and the addition of the cryIIIA transcription terminator at the 3' end of the lacZ gene, produced only marginal effects on the final beta-galactosidase activity.

L6 ANSWER 6 OF 6 HCAPLUS COPYRIGHT 2006 ACS on STN  
ACCESSION NUMBER: 1999:566201 HCAPLUS  
DOCUMENT NUMBER: 131:180803  
TITLE: Nucleic acid vectors for recombinant production of heterologous proteins in a **Bacillus** cell  
INVENTOR(S): Widner, William; Sloma, Alan; Thomas, Michael D.  
PATENT ASSIGNEE(S): Novo Nordisk Biotech, Inc., USA  
SOURCE: PCT Int. Appl., 90 pp.  
CODEN: PIXXD2  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
FAMILY ACC. NUM. COUNT: 3  
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9943835	A2	19990902	WO 1999-US4360	19990226
WO 9943835	A3	19991125		

W: AL, AU, BB, BG, BR, CA, CN, CU, CZ, EE, GE, HU, IL, IN, IS, JP, KP, KR, LC, LK, LR, LT, LU, LV, MG, MK, MN, MX, NO, NZ, PL, RO, SG, SI, SK, TR, TT, UA, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM

RW: GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG

US 5955310	A	19990921	US 1998-31442	19980226
AU 9929756	A1	19990915	AU 1999-29756	19990226
EP 1056873	A2	20001206	EP 1999-911012	19990226
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, NL, SE, PT, IE, FI				
JP 2002504379	T2	20020212	JP 2000-533574	19990226
CN 1510145	A	20040707	CN 2003-2003158121	19990226
US 2003170876	A1	20030911	US 2001-834271	20010412
PRIORITY APPLN. INFO.:			US 1998-31442	A 19980226
			US 1999-256377	B3 19990224
			WO 1999-US4360	W 19990226

AB The present invention relates to methods for producing a polypeptide, comprising: (a) cultivating a *Bacillus* host cell in a medium conducive for the production of the polypeptide, wherein the *Bacillus* cell comprises a nucleic acid construct comprising (i) a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and alternatively also (ii) an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide; and (b) isolating the polypeptide from the cultivation medium. The present invention also relates to methods for producing a polypeptide, comprising: (a) cultivating a *Bacillus* host cell in a medium conducive for the production of the polypeptide, wherein the *Bacillus* cell comprises a nucleic acid construct comprising (i) a "consensus" promoter having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide and (ii) an mRNA processing/stabilizing sequence located downstream of the "consensus" promoter and upstream of the nucleic acid sequence encoding the polypeptide; and (b) isolating the polypeptide from the cultivation medium. Random promoters are created by placing promoters such as amyQ and amyL upstream of the cryIIIA promoter and its mRNA stabilizing sequence. Alternatively, "consensus" amyQ promoters are created with the cryIIIA mRNA stabilizing sequence, as well as tandem copies of a single promoter such as the short consensus amyQ dimer and trimer promoters. All of these approaches lead to significantly higher levels of SAVINASE gene expression (up to 620%) in *Bacillus* cells when compared to the levels obtained using single promoters such as amyQ and amyL.

=> d his

(FILE 'HOME' ENTERED AT 14:23:15 ON 23 FEB 2006)

FILE 'MEDLINE, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH, HCAPLUS, NTIS, LIFESCI' ENTERED AT 14:23:47 ON 23 FEB 2006

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L1      395086 S BACILLUS
L2      560 S CRYIIIA
L3      516 S L1 AND L2
L4      635 S "TTGACA" OR "TATAAT"
L5      13 S L3 AND L4
L6      6 DUP REM L5 (7 DUPLICATES REMOVED)
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=> s l2 and "consensus promoter"

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L7      7 L2 AND "CONSENSUS PROMOTER"
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=> dup rem l7

PROCESSING COMPLETED FOR L7

L8 5 DUP REM L7 (2 DUPLICATES REMOVED)

=> d 1-5 ibib ab

L8 ANSWER 1 OF 5 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2005-30339 BIOTECHDS

TITLE: Producing a hyaluronic acid, by cultivating Bacillus cell comprising a nucleic acid construct comprising a variant amyL promoter, a **consensus promoter**, and a **cryIIIA** promoter, and isolating hyaluronic acid from the cultivation medium;  
production of recombinant hyaluronic acid from a Bacillus having a triple promoter useful for a tissue engineering application

AUTHOR: WIDNER W; SLOMA A; THOMAS M; TANG M

PATENT ASSIGNEE: NOVOZYMES BIOPOLYMER AS

PATENT INFO: US 2005221446 6 Oct 2005

APPLICATION INFO: US 2005-96190 31 Mar 2005

PRIORITY INFO: US 2005-96190 31 Mar 2005; US 2004-558507 31 Mar 2004

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: WPI: 2005-722702 [74]

AB DERWENT ABSTRACT:

NOVELTY - Producing a hyaluronic acid comprises cultivating a Bacillus cell in a medium for the production of the hyaluronic acid, where the Bacillus cell comprises a nucleic acid construct comprising a triple promoter comprising a variant amyL promoter, a **consensus promoter**, and a **cryIIIA** promoter, and isolating the hyaluronic acid from the cultivation medium.

DETAILED DESCRIPTION - Producing a hyaluronic acid comprises: (A) cultivating a Bacillus cell in a medium for the production of the hyaluronic acid, where the Bacillus cell comprises a nucleic acid construct comprising a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of a fully defined 614 bp sequence (SEQ ID NO. 1), a **consensus promoter** having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region, and a **cryIIIA** promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of the hyaluronic acid; and (B) isolating the hyaluronic acid from the cultivation medium. INDEPENDENT CLAIMS are also included for: (1) a Bacillus cell comprising a nucleic acid construct which comprises: (a) a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of SEQ ID NO. 1, a **consensus promoter** having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region, and a **cryIIIA** promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of a hyaluronic acid, and optionally (b) an mRNA processing/stabilizing sequence located downstream of the triple promoter and upstream of the one or more coding sequences involved in the biosynthesis of the hyaluronic acid; (2) producing a selectable marker-free mutant of a Bacillus cell; and (3) a selectable marker-free mutant of a Bacillus cell obtained by the method above.

WIDER DISCLOSURE - Also disclosed are: (1) methods for obtaining a Bacillus host cell; and (2) nucleic acid constructs comprising a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of SEQ ID NO. 1, a **consensus promoter** having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region, and a **cryIIIA** promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of a hyaluronic acid.

BIOTECHNOLOGY - Preferred Method: In producing a hyaluronic acid,

the variant amyL promoter is SEQ ID NO. 1. The consensus promoter is obtained from the Bacillus amyloliquefaciens alpha-amylase gene (amyQ), where the consensus amyQ promoter has the nucleotide sequence comprising fully defined 185 bp sequence (SEQ ID NO. 42 or 43). The nucleic acid construct further comprises an mRNA processing/stabilizing sequence located downstream of the triple promoter and upstream of the one or more coding sequences involved in the biosynthesis of the hyaluronic acid. Preferably, the one or more coding sequences involved in the biosynthesis of the hyaluronic acid are selected from a hyaluronan synthase, UDP-glucose 6-dehydrogenase, UDP-glucose pyrophosphorylase, UDP-N-acetylglucosamine pyrophosphorylase, glucose-6-phosphate isomerase, hexokinase, phosphoglucomutase, amidotransferase, mutase, or acetyl transferase gene. Producing a selectable marker-free mutant of a Bacillus cell comprises deleting a selectable marker gene of the Bacillus cell. The Bacillus cell contains no foreign selectable marker gene.

USE - The method is useful for producing a hyaluronic acid. Hyaluronic acid is useful in eye and joint surgery. Products of hyaluronic acid are also useful in orthopedics, rheumatology, and in dermatology.

EXAMPLE - No relevant example given. (94 pages)

L8 ANSWER 2 OF 5 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN  
DUPLICATE 1

ACCESSION NUMBER: 2004-02033 BIOTECHDS

TITLE: Generating an expression library of polynucleotides by introducing the linear integration cassette into the host cell and selecting or screening for host cells that produce the polypeptides of interest;  
protein library screening using homologous recombination

AUTHOR: BJORNVAAD M E; JORGENSEN P L; HANSEN P K

PATENT ASSIGNEE: NOVOZYMES AS

PATENT INFO: WO 2003095658 20 Nov 2003

APPLICATION INFO: WO 2003-DK301 7 May 2003

PRIORITY INFO: DK 2002-682 7 May 2002; DK 2002-682 7 May 2002

DOCUMENT TYPE: Patent

LANGUAGE: English

AB DERWENT ABSTRACT:

NOVELTY - Generating an expression library of polynucleotides integrated by homologous recombination into the genome of a competent Gram-positive bacterium host cell comprises: (a) providing a non-replicating linear integration cassette; and (b) introducing the linear integration cassette into the host cell and selecting or screening for host cells that produce the polypeptides of interest.

DETAILED DESCRIPTION - Generating an expression library of polynucleotides integrated by homologous recombination into the genome of a competent Gram-positive bacterium host cell comprises: (a) providing a non-replicating linear integration cassette; and (b) introducing the linear integration cassette into the host cell and selecting or screening for host cells that produce the polypeptides of interest. The cassette comprises: (1) a polynucleotide encoding one or more polypeptides of interest; (2) a 5' flanking polynucleotide segment upstream of the polynucleotide of (1) and comprising a first homologous region located in the 3' end of the segment; and (3) a 3' flanking polynucleotide segment downstream of the polynucleotide of (1) and comprising a second homologous region located in the 5' end of the segment. The first and second homologous regions are at least 500, 1000 or 1500 bp, each of which has a sequence identity of at least 80, 85, 90 or 95-100% with a region of the host cell genom. INDEPENDENT CLAIMS are also included for the following: (1) a non-replicating linear Gram-positive host cell integration cassette; and (2) a method of producing a polypeptide of interest.

BIOTECHNOLOGY - Preferred Method: Generating an expression library of polynucleotides integrated by homologous recombination into the genome

of a competent Gram-positive bacterium host cell comprises the additional step carried out between steps (1) and (2) that comprises introducing the plasmid into an intermediate *Escherichia coli* host cell and propagating it by replication. The integration cassette comprises: (1) an mRNA processing/stabilizing sequence derived from *cryIIIA*-gene; (2) a terminator downstream of the polynucleotide encoding the polypeptides of interest; and (3) a multiple cloning site with at least one recognition site for a restriction nuclease. It further comprises a marker gene located between the 5' and 3' flanking segments and at least one promoter that is a consensus promoter having the sequence TTGACA for the -35 region and TATAAT for the -10 region, and/or which is derived from *amyL*, *amyQ*, *amyM*, *cryIIIA*, *dagA*, *aprH*, *penP*, *sacB*, *spol*, *tac*, *xylA* or *xylB*. The promoter is located between the flanking segments and is operably linked to the polynucleotide encoding one or more polypeptides of interest. Each of the 5' and 3' flanking polynucleotide segments comprises at least 500, 1000, 1500 or 2000 bp of non-homologous polynucleotides located in the 5' and 3' end of the 5' and 3' flanking segments, respectively. The promoter is one that results in that the host cells produce the polypeptides of interest in a yield of at least 10 mg/L. The polynucleotide comprises natural, synthetic or a library of shuffled or recombined homologs or variants of a gene or operon, provided by DNA breeding or DNA shuffling. The polypeptides of interest comprise enzymes, proteins or antimicrobial peptides. The enzymes are involved in the biosynthesis of hyaluronic acid. The Gram-positive host cell is *Bacillus subtilis*. The homologous region of the 5' and/or the 3' flanking segment is comprised in the *yfmD*-*yfmC*-*yfmB*-*yfmA*-*pelB*-*yflS*-*cits* region of the *Bacillus subtilis* genome or in the *cryIIIA* promoter. The non-replicating linear integration cassette is comprised in a plasmid and introduced into the host cell. The plasmid is capable of replicating in an *Escherichia coli* host cell but not in a *Bacillus* host cell. Producing a polypeptide of interest comprises culturing Gram-positive host cells comprising the integration cassette integrated into its genome, under conditions promoting expression of the polypeptide of interest. The method further comprises isolating and/or purifying the polypeptide of interest.

USE - The method is useful in generating an expression library of polynucleotides integrated by homologous recombination into the genome of a competent Gram-positive bacterium host cell (claimed). (55 pages)

L8 ANSWER 3 OF 5 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN  
DUPLICATE 2

ACCESSION NUMBER: 2004-08483 BIOTECHDS

TITLE: Production of a secreted polypeptide having L-asparaginase activity for treating leukemia, comprises cultivating a host cell comprising a nucleic acid having a sequence encoding a secretory signal peptide linked to a second sequence;  
vector-mediated enzyme gene transfer and expression in host cell for recombinant protein production, amino acid preparation and disease therapy

AUTHOR: THOMAS M D; SLOMA A

PATENT ASSIGNEE: NOVOZYMES BIOTECH INC

PATENT INFO: US 2003186380 2 Oct 2003

APPLICATION INFO: US 2003-406025 1 Apr 2003

PRIORITY INFO: US 2003-406025 1 Apr 2003; US 2002-369192 1 Apr 2002

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: WPI: 2004-088916 [09]

AB DERWENT ABSTRACT:

NOVELTY - Producing a secreted polypeptide having L-asparaginase activity, comprises cultivating a host cell containing a nucleic acid construct having a sequence encoding a secretory signal peptide linked to a second sequence encoding the polypeptide having L-asparaginase activity, where the signal peptide directs the polypeptide into the cell's secretory pathway, and recovering the secreted polypeptide.

DETAILED DESCRIPTION - An INDEPENDENT CLAIM is also included for a recombinant expression vector comprising the nucleic acid construct.

BIOTECHNOLOGY - Preferred Material: The polypeptide is encoded by a nucleic acid sequence contained in plasmid pCR2.1-yccC, which is contained in *Escherichia coli* (NRRL B-30558). The consensus promoter is obtained from a promoter obtained from the *E. coli* lac operon *Streptomyces coelicolor* agarase gene (*dagA*), *Bacillus clausii* alkaline protease gene (*aprH*), *B. licheniformis* alkaline protease gene (*subtilisin* Carlsberg gene), *B. subtilis* levansucrase gene (*sacB*), *B. subtilis* alpha-amylase gene (*amyE*), *B. licheniformis* alpha-amylase gene (*amyL*), *B. stearothermophilus* maltogenic amylase gene (*amyM*), *B. licheniformis* penicillinase gene (*penP*), *B. subtilis* *xylA* and *xylB* genes, *B. thuringiensis* subsp. *tenebrionis* *CryIIIA* gene (*cryIIIA*) or its portions, or preferably *B. amyloliquefaciens* alpha-amylase gene (*amyQ*). The mRNA processing/stabilizing sequence is the *cryIIIA* mRNA processing/stabilizing sequence. The bacillus cell is *B. alkalophilus*, *B. amyloliquefaciens*, *B. brevis*, *B. circulans*, *B. clausii*, *B. coagulans*, *B. lautus*, *B. lentus*, *B. licheniformis*, *B. megaterium*, *B. stearothermophilus*, *B. subtilis*, or *B. thuringiensis*. Preferred Component: The nucleic acid construct comprises a tandem promoter, in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding a polypeptide, and optionally an mRNA processing stabilizing sequence located downstream of the tandem promoter and upstream of the second nucleic acid sequence, encoding the polypeptide having L-asparaginase activity. It comprises a consensus promoter having the sequence TTGACA for the 35 region, and TATAAT for the 10 region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and an mRNA processing stabilizing sequence located downstream of the consensus promoter, and upstream of the second nucleic acid sequence encoding the polypeptide having L-asparagine activity. The consensus promoter is obtained from any bacterial or a bacillus promoter. The nucleic acid construct comprises a ribosome binding site sequence heterologous to the host cell.

ACTIVITY - Cytostatic.

MECHANISM OF ACTION - L-asparaginase.

USE - The invention is for production of secreted polypeptide having L-asparaginase activity, for use in producing L-aspartate from L-asparagine. The inventive secreted polypeptide is useful for treatment of leukemia, e.g. acute lymphocytic leukemia.

ADVANTAGE - The invention achieves secretion of L-asparaginase enabling easy recovery and purification, high expression constructs for producing the L-asparaginase in high amounts, and the use of host cells for production that have generally regarded as safe status.

EXAMPLE - *B. subtilis* strains MDT51 and MDT52 were grown in Lactobacilli MRS Broth (RTM; 50 ml) at 37degreesC, and 250 revolutions/minute (rpm) for 24 hours. Supernatants were recovered by centrifugation at 7000 rpm for 5 minutes. A prominent band corresponding to a protein of the expected size for mature L-asparagine (37 kDa; amino acids 24-375) was observed in the MDT51 sample, but not in the MDT52 sample. (22 pages)

L8 ANSWER 4 OF 5 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-04169 BIOTECHDS

TITLE: Producing a polypeptide comprises cultivating a *Bacillus* cell in a medium conducive to the production of the polypeptide, where the *Bacillus* cell comprises a nucleic acid construct comprising a tandem promoter;  
involving vector-mediated gene transfer and expression in host cell for use as a selectable marker

AUTHOR: WIDNER W; SLOMA A; THOMAS M D

PATENT ASSIGNEE: NOVOZYMES BIOTECH INC

PATENT INFO: US 2003170876 11 Sep 2003

APPLICATION INFO: US 2001-834271 12 Apr 2001  
PRIORITY INFO: US 2001-834271 12 Apr 2001; US 1998-31442 26 Feb 1998  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
OTHER SOURCE: WPI: 2003-898275 [82]  
AB DERWENT ABSTRACT:

NOVELTY - Producing a polypeptide comprises cultivating a *Bacillus* cell in a medium conducive to the production of the polypeptide, where the *Bacillus* cell comprises a nucleic acid construct comprising a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a nucleic acid sequence encoding the polypeptide, and isolating the polypeptide from the cultivation medium.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are included for the following: (1) a *Bacillus* cell comprising a nucleic acid construct comprising a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding a polypeptide, and optionally an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide; (2) a method for obtaining a *Bacillus* host cell by introducing into a *Bacillus* cell the nucleic acid construct cited above; (3) a method for producing a selectable marker-free mutant of a *Bacillus* cell by deleting a selectable marker gene of the *Bacillus* cell; and (4) a selectable marker-free mutant of a *Bacillus* cell.

BIOTECHNOLOGY - Preferred Method: In producing a polypeptide, the nucleic acid construct further comprises an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide. The tandem promoter comprises two or more bacterial promoter sequences, which are obtained from one or more *Bacillus* genes. The tandem promoter comprises the amyQ promoter, a consensus promoter having the sequence TTGACA for the -35 region and TATAAT from the -10 region, the amyL promoter, and/or the cryIIIA promoter. The tandem promoter comprises two copies of the amyQ, amyL or cryIIIA promoter. The two or more promoter sequences of the tandem promoter simultaneously promote the transcription of the nucleic acid sequence. The one or more of the two or more promoter sequences of the tandem promoter promote the transcription of the nucleic acid sequence at different stages of growth of the *Bacillus* cell. The mRNA processing/stabilizing sequence is the cryIIIA or SP82 mRNA processing/stabilizing sequence, which generates mRNA transcripts essentially of the same size. The *Bacillus* cell contains one or more copies of the nucleic acid construct. The nucleic acid construct further comprises a selectable marker gene. The nucleic acid sequence encodes a polypeptide heterologous to the *Bacillus* cell. The polypeptide is a hormone or its variant, enzyme, receptor or its portion, antibody or its portion, or reporter. The enzyme is an oxidoreductase, transferase, hydrolase, lyase, isomerase or ligase. The enzyme is preferably an aminopeptidase, amylase, carbohydrase, carboxypeptidase, catalase, cellulase, chitinase, cutinase, cyclodextrin glycosyltransferase, deoxyribonuclease, esterase, alpha-galactosidase, beta-galactosidase, glucoamylase, alpha-glucosidase, beta-glucosidase, invertase, laccase, lipase, mannosidase, mutanase, oxidase, a pectinolytic enzyme, peroxidase, phytase, polyphenoloxidase, proteolytic enzyme, ribonuclease, transglutaminase or xylanase. The nucleic acid sequence is contained in the chromosome of the *Bacillus* cell. The *Bacillus* host cell is *Bacillus alkalophilus*, *Bacillus amyloliquefaciens*, *Bacillus brevis*, *Bacillus brevis*, *Bacillus circulans*, *Bacillus clausii*, *Bacillus coagulans*, *Bacillus firmus*, *Bacillus lautus*, *Bacillus lentus*, *Bacillus licheniformis*, *Bacillus megaterium*, *Bacillus pumilus*, *Bacillus sterothermophilus*, *Bacillus subtilis*, or *Bacillus thuringiensis*. This method alternatively comprises cultivating a *Bacillus* cell in a medium conducive for the production of the polypeptide, where the *Bacillus* cell comprises a nucleic acid construct comprising a consensus promoter having the sequence TTGACA for the -35 region and TATAAT

for the -10 region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and an mRNA processing/stabilizing sequence located downstream of the consensus promoter and upstream of the nucleic acid sequence encoding the polypeptide; and isolating the polypeptide from the cultivation medium. The consensus promoter is obtained from any bacterial promoter, preferably a Bacillus promoter. Preferred Cell: The Bacillus cell comprises a nucleic acid construct that further comprises a selectable marker gene. The cell can also contain no selectable marker gene.

USE - The methods are useful for producing a polypeptide in a Bacillus cell, and for producing a selectable marker-free mutant of a Bacillus cell.

EXAMPLE - No relevant example given. (57 pages)

L8 ANSWER 5 OF 5 HCAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 1999:566201 HCAPLUS  
DOCUMENT NUMBER: 131:180803  
TITLE: Nucleic acid vectors for recombinant production of heterologous proteins in a Bacillus cell  
INVENTOR(S): Widner, William; Sloma, Alan; Thomas, Michael D.  
PATENT ASSIGNEE(S): Novo Nordisk Biotech, Inc., USA  
SOURCE: PCT Int. Appl., 90 pp.  
CODEN: PIXXD2  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
FAMILY ACC. NUM. COUNT: 3  
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 9943835	A2	19990902	WO 1999-US4360	19990226
WO 9943835	A3	19991125		
W:	AL, AU, BB, BG, BR, CA, CN, CU, CZ, EE, GE, HU, IL, IN, IS, JP, KP, KR, LC, LK, LR, LT, LU, LV, MG, MK, MN, MX, NO, NZ, PL, RO, SG, SI, SK, TR, TT, UA, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
RW:	GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG			
US 5955310	A	19990921	US 1998-31442	19980226
AU 9929756	A1	19990915	AU 1999-29756	19990226
EP 1056873	A2	20001206	EP 1999-911012	19990226
R:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, NL, SE, PT, IE, FI			
JP 2002504379	T2	20020212	JP 2000-533574	19990226
CN 1510145	A	20040707	CN 2003-2003158121	19990226
US 2003170876	A1	20030911	US 2001-834271	20010412
PRIORITY APPLN. INFO.:			US 1998-31442	A 19980226
			US 1999-256377	B3 19990224
			WO 1999-US4360	W 19990226

AB The present invention relates to methods for producing a polypeptide, comprising: (a) cultivating a Bacillus host cell in a medium conducive for the production of the polypeptide, wherein the Bacillus cell comprises a nucleic acid construct comprising (i) a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and alternatively also (ii) an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide; and (b) isolating the polypeptide from the cultivation medium. The present invention also relates to methods for producing a polypeptide, comprising: (a) cultivating a Bacillus host cell in a medium conducive for the production of the polypeptide, wherein the Bacillus cell comprises a nucleic acid construct comprising (i) a "consensus" promoter having the sequence TTGACA for the



"-35" region and TATAAT for the "-10" region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide and (ii) an mRNA processing/stabilizing sequence located downstream of the "consensus" promoter and upstream of the nucleic acid sequence encoding the polypeptide; and (b) isolating the polypeptide from the cultivation medium. Random promoters are created by placing promoters such as amyQ and amyL upstream of the cryIIIA promoter and its mRNA stabilizing sequence. Alternatively, "consensus" amyQ promoters are created with the cryIIIA mRNA stabilizing sequence, as well as tandem copies of a single promoter such as the short consensus amyQ dimer and trimer promoters. All of these approaches lead to significantly higher levels of SAVINASE gene expression (up to 620%) in Bacillus cells when compared to the levels obtained using single promoters such as amyQ and amyL.

=> d his

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FILE 'MEDLINE, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH, HCAPLUS, NTIS, LIFESCI' ENTERED AT 14:23:47 ON 23 FEB 2006

```
L1      395086 S BACILLUS
L2      560 S CRYIIIA
L3      516 S L1 AND L2
L4      635 S "TTGACA" OR "TATAAT"
L5      13 S L3 AND L4
L6      6 DUP REM L5 (7 DUPLICATES REMOVED)
L7      7 S L2 AND "CONSENSUS PROMOTER"
L8      5 DUP REM L7 (2 DUPLICATES REMOVED)
```

=> e widner w/au

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E1      32      WIDNER T E/AU
E2      2      WIDNER THOMAS E/AU
E3      31 --> WIDNER W/AU
E4      7      WIDNER W E/AU
E5      43      WIDNER W R/AU
E6      14      WIDNER WILLIAM/AU
E7      22      WIDNER WILLIAM R/AU
E8      1      WIDNER WILLIAM ROY/AU
E9      1      WIDNER WM R/AU
E10     5      WIDNES C/AU
E11     2      WIDNES J/AU
E12     2      WIDNES J A/AU
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=> s e3-e9

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L9      119 ("WIDNER W"/AU OR "WIDNER W E"/AU OR "WIDNER W R"/AU OR "WIDNER
WILLIAM"/AU OR "WIDNER WILLIAM R"/AU OR "WIDNER WILLIAM ROY"/AU
OR "WIDNER WM R"/AU)
```

=> e sloma a/au

```
E1      3      SLOM T J/AU
E2      2      SLOM TREVOR J/AU
E3      124 --> SLOMA A/AU
E4      15      SLOMA A P/AU
E5      60      SLOMA ALAN/AU
E6      16      SLOMA ALAN P/AU
E7      1      SLOMA ALAN PAUL/AU
E8      5      SLOMA C R/AU
E9      4      SLOMA CARI R/AU
E10     5      SLOMA CARI ROARK/AU
E11     1      SLOMA D/AU
E12     1      SLOMA D R/AU
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=> s e3  
L10 124 "SLOMA A"/AU

=> e thomas m d/au  
E1 16 THOMAS M CARMEN/AU  
E2 1 THOMAS M CUARTERO C/AU  
E3 404 --> THOMAS M D/AU  
E4 105 THOMAS M D A/AU  
E5 4 THOMAS M D H/AU  
E6 3 THOMAS M D JR/AU  
E7 2 THOMAS M D O/AU  
E8 17 THOMAS M D R/AU  
E9 346 THOMAS M E/AU  
E10 33 THOMAS M E A/AU  
E11 5 THOMAS M E C/AU  
E12 31 THOMAS M E M/AU

=> s e3  
L11 404 "THOMAS M D"/AU

=> d his

(FILE 'HOME' ENTERED AT 14:23:15 ON 23 FEB 2006)

FILE 'MEDLINE, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH, HCAPLUS, NTIS, LIFESCI' ENTERED AT 14:23:47 ON 23 FEB 2006

L1 395086 S BACILLUS  
L2 560 S CRYIIIA  
L3 516 S L1 AND L2  
L4 635 S "TTGACA" OR "TATAAT"  
L5 13 S L3 AND L4  
L6 6 DUP REM L5 (7 DUPLICATES REMOVED)  
L7 7 S L2 AND "CONSENSUS PROMOTER"  
L8 5 DUP REM L7 (2 DUPLICATES REMOVED)  
E WIDNER W/AU  
L9 119 S E3-E9  
E SLOMA A/AU  
L10 124 S E3  
E THOMAS M D/AU  
L11 404 S E3

=> s l9 or l10 or l11  
L12 632 L9 OR L10 OR L11

=> s l2 and l12  
L13 8 L2 AND L12

=> dup rem l13  
PROCESSING COMPLETED FOR L13  
L14 6 DUP REM L13 (2 DUPLICATES REMOVED)

=> d 1-6 ibib ab

L14 ANSWER 1 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN  
DUPLICATE 1

ACCESSION NUMBER: 2005-30339 BIOTECHDS

TITLE: Producing a hyaluronic acid, by cultivating Bacillus cell comprising a nucleic acid construct comprising a variant amyL promoter, a consensus promoter, and a cryIIIA promoter, and isolating hyaluronic acid from the cultivation medium;  
production of recombinant hyaluronic acid from a Bacillus having a triple promoter useful for a tissue engineering application

AUTHOR: WIDNER W; SLOMA A; THOMAS M; TANG M  
PATENT ASSIGNEE: NOVOZYMES BIOPOLYMER AS  
PATENT INFO: US 2005221446 6 Oct 2005  
APPLICATION INFO: US 2005-96190 31 Mar 2005  
PRIORITY INFO: US 2005-96190 31 Mar 2005; US 2004-558507 31 Mar 2004  
DOCUMENT TYPE: Patent  
LANGUAGE: English  
OTHER SOURCE: WPI: 2005-722702 [74]

AB DERWENT ABSTRACT:

NOVELTY - Producing a hyaluronic acid comprises cultivating a *Bacillus* cell in a medium for the production of the hyaluronic acid, where the *Bacillus* cell comprises a nucleic acid construct comprising a triple promoter comprising a variant amyL promoter, a consensus promoter, and a cryIIIA promoter, and isolating the hyaluronic acid from the cultivation medium.

DETAILED DESCRIPTION - Producing a hyaluronic acid comprises: (A) cultivating a *Bacillus* cell in a medium for the production of the hyaluronic acid, where the *Bacillus* cell comprises a nucleic acid construct comprising a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of a fully defined 614 bp sequence (SEQ ID NO. 1), a consensus promoter having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region, and a cryIIIA promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of the hyaluronic acid; and (B) isolating the hyaluronic acid from the cultivation medium. INDEPENDENT CLAIMS are also included for: (1) a *Bacillus* cell comprising a nucleic acid construct which comprises: (a) a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of SEQ ID NO. 1, a consensus promoter having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region, and a cryIIIA promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of a hyaluronic acid, and optionally (b) an mRNA processing/stabilizing sequence located downstream of the triple promoter and upstream of the one or more coding sequences involved in the biosynthesis of the hyaluronic acid; (2) producing a selectable marker-free mutant of a *Bacillus* cell; and (3) a selectable marker-free mutant of a *Bacillus* cell obtained by the method above.

WIDER DISCLOSURE - Also disclosed are: (1) methods for obtaining a *Bacillus* host cell; and (2) nucleic acid constructs comprising a triple promoter comprising a variant amyL promoter having a mutation corresponding to position 590 of SEQ ID NO. 1, a consensus promoter having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region, and a cryIIIA promoter, in which each promoter sequence of the triple promoter is operably linked to one or more coding sequences involved in the biosynthesis of a hyaluronic acid.

BIOTECHNOLOGY - Preferred Method: In producing a hyaluronic acid, the variant amyL promoter is SEQ ID NO. 1. The consensus promoter is obtained from the *Bacillus amyloliquefaciens* alpha-amylase gene (amyQ), where the consensus amyQ promoter has the nucleotide sequence comprising fully defined 185 bp sequence (SEQ ID NO. 42 or 43). The nucleic acid construct further comprises an mRNA processing/stabilizing sequence located downstream of the triple promoter and upstream of the one or more coding sequences involved in the biosynthesis of the hyaluronic acid. Preferably, the one or more coding sequences involved in the biosynthesis of the hyaluronic acid are selected from a hyaluronan synthase, UDP-glucose 6-dehydrogenase, UDP-glucose pyrophosphorylase, UDP-N-acetylglucosamine pyrophosphorylase, glucose-6-phosphate isomerase, hexokinase, phosphoglucomutase, amidotransferase, mutase, or acetyl transferase gene. Producing a selectable marker-free mutant of a *Bacillus* cell comprises deleting a selectable marker gene of the *Bacillus* cell. The *Bacillus* cell contains no foreign selectable marker gene.

USE - The method is useful for producing a hyaluronic acid.

Hyaluronic acid is useful in eye and joint surgery. Products of hyaluronic acid are also useful in orthopedics, rheumatology, and in dermatology.

EXAMPLE - No relevant example given. (94 pages)

L14 ANSWER 2 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-08483 BIOTECHDS

TITLE: Production of a secreted polypeptide having L-asparaginase activity for treating leukemia, comprises cultivating a host cell comprising a nucleic acid having a sequence encoding a secretory signal peptide linked to a second sequence; vector-mediated enzyme gene transfer and expression in host cell for recombinant protein production, amino acid preparation and disease therapy

AUTHOR: THOMAS M D; SLOMA A

PATENT ASSIGNEE: NOVOZYMES BIOTECH INC

PATENT INFO: US 2003186380 2 Oct 2003

APPLICATION INFO: US 2003-406025 1 Apr 2003

PRIORITY INFO: US 2003-406025 1 Apr 2003; US 2002-369192 1 Apr 2002

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: WPI: 2004-088916 [09]

AB DERWENT ABSTRACT:

NOVELTY - Producing a secreted polypeptide having L-asparaginase activity, comprises cultivating a host cell containing a nucleic acid construct having a sequence encoding a secretory signal peptide linked to a second sequence encoding the polypeptide having L-asparaginase activity, where the signal peptide directs the polypeptide into the cell's secretory pathway, and recovering the secreted polypeptide.

DETAILED DESCRIPTION - An INDEPENDENT CLAIM is also included for a recombinant expression vector comprising the nucleic acid construct.

BIOTECHNOLOGY - Preferred Material: The polypeptide is encoded by a nucleic acid sequence contained in plasmid pCR2.1-yccc, which is contained in *Escherichia coli* (NRRL B-30558). The consensus promoter is obtained from a promoter obtained from the *E. coli* lac operon *Streptomyces coelicolor* agarase gene (*dagA*), *Bacillus clausii* alkaline protease gene (*aprH*), *B. licheniformis* alkaline protease gene (*subtilisin* Carlsberg gene), *B. subtilis* levansucrase gene (*sacB*), *B. subtilis* alpha-amylase gene (*amyE*), *B. licheniformis* alpha-amylase gene (*amyL*), *B. stearothermophilus* maltogenic amylase gene (*amyM*), *B. licheniformis* penicillinase gene (*penP*), *B. subtilis* *xylA* and *xylB* genes, *B. thuringiensis* subsp. *tenebrionis* *CryIIIA* gene (*cryIIIA*) or its portions, or preferably *B. amyloliquefaciens* alpha-amylase gene (*amyQ*). The mRNA processing/stabilizing sequence is the *cryIIIA* mRNA processing/stabilizing sequence. The bacillus cell is *B. alkalophilus*, *B. amyloliquefaciens*, *B. brevis*, *B. circulans*, *B. clausii*, *B. coagulans*, *B. lautus*, *B. lentus*, *B. licheniformis*, *B. megaterium*, *B. stearothermophilus*, *B. subtilis*, or *B. thuringiensis*. Preferred

Component: The nucleic acid construct comprises a tandem promoter, in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding a polypeptide, and optionally an mRNA processing stabilizing sequence located downstream of the tandem promoter and upstream of the second nucleic acid sequence, encoding the polypeptide having L-asparaginase activity. It comprises a consensus promoter having the sequence TTGACA for the 35 region, and TATAAT for the 10 region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and an mRNA processing stabilizing sequence located downstream of the consensus promoter, and upstream of the second nucleic acid sequence encoding the polypeptide having L-asparagine activity. The consensus promoter is obtained from any bacterial or a bacillus promoter. The nucleic acid construct comprises a ribosome binding site sequence heterologous to the host cell.

ACTIVITY - Cytostatic.

MECHANISM OF ACTION - L-asparaginase.

USE - The invention is for production of secreted polypeptide having L-asparaginase activity, for use in producing L-aspartate from L-asparagine. The inventive secreted polypeptide is useful for treatment of leukemia, e.g. acute lymphocytic leukemia.

ADVANTAGE - The invention achieves secretion of L-asparaginase enabling easy recovery and purification, high expression constructs for producing the L-asparaginase in high amounts, and the use of host cells for production that have generally regarded as safe status.

EXAMPLE - *B. subtilis* strains MDT51 and MDT52 were grown in Lactobacilli MRS Broth (RTM; 50 ml) at 37degreesC, and 250 revolutions/minute (rpm) for 24 hours. Supernatants were recovered by centrifugation at 7000 rpm for 5 minutes. A prominent band corresponding to a protein of the expected size for mature L-asparagine (37 kDa; amino acids 24-375) was observed in the MDT51 sample, but not in the MDT52 sample. (22 pages)

L14 ANSWER 3 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN

ACCESSION NUMBER: 2004-04169 BIOTECHDS

TITLE: Producing a polypeptide comprises cultivating a *Bacillus* cell in a medium conducive to the production of the polypeptide, where the *Bacillus* cell comprises a nucleic acid construct comprising a tandem promoter;  
involving vector-mediated gene transfer and expression in host cell for use as a selectable marker

AUTHOR: WIDNER W; SLOMA A; THOMAS M D

PATENT ASSIGNEE: NOVOZYMES BIOTECH INC

PATENT INFO: US 2003170876 11 Sep 2003

APPLICATION INFO: US 2001-834271 12 Apr 2001

PRIORITY INFO: US 2001-834271 12 Apr 2001; US 1998-31442 26 Feb 1998

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: WPI: 2003-898275 [82]

AB DERWENT ABSTRACT:

NOVELTY - Producing a polypeptide comprises cultivating a *Bacillus* cell in a medium conducive to the production of the polypeptide, where the *Bacillus* cell comprises a nucleic acid construct comprising a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a nucleic acid sequence encoding the polypeptide, and isolating the polypeptide from the cultivation medium.

DETAILED DESCRIPTION - INDEPENDENT CLAIMS are included for the following: (1) a *Bacillus* cell comprising a nucleic acid construct comprising a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding a polypeptide, and optionally an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide; (2) a method for obtaining a *Bacillus* host cell by introducing into a *Bacillus* cell the nucleic acid construct cited above; (3) a method for producing a selectable marker-free mutant of a *Bacillus* cell by deleting a selectable marker gene of the *Bacillus* cell; and (4) a selectable marker-free mutant of a *Bacillus* cell.

BIOTECHNOLOGY - Preferred Method: In producing a polypeptide, the nucleic acid construct further comprises an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide. The tandem promoter comprises two or more bacterial promoter sequences, which are obtained from one or more *Bacillus* genes. The tandem promoter comprises the amyQ promoter, a consensus promoter having the sequence TTGACA for the -35 region and TATAAT from the -10 region, the amyL promoter, and/or the cryIIIA promoter. The tandem promoter comprises two copies of the amyQ, amyL or cryIIIA promoter. The two or more promoter sequences of the tandem promoter simultaneously promote the transcription of the nucleic acid sequence. The one or more of the two or more promoter sequences of the tandem promoter promote the transcription of the nucleic

acid sequence at different stages of growth of the *Bacillus* cell. The mRNA processing/stabilizing sequence is the *cryIIIA* or SP82 mRNA processing/stabilizing sequence, which generates mRNA transcripts essentially of the same size. The *Bacillus* cell contains one or more copies of the nucleic acid construct. The nucleic acid construct further comprises a selectable marker gene. The nucleic acid sequence encodes a polypeptide heterologous to the *Bacillus* cell. The polypeptide is a hormone or its variant, enzyme, receptor or its portion, antibody or its portion, or reporter. The enzyme is an oxidoreductase, transferase, hydrolase, lyase, isomerase or ligase. The enzyme is preferably an aminopeptidase, amylase, carbohydrazase, carboxypeptidase, catalase, cellulase, chitinase, cutinase, cyclodextrin glycosyltransferase, deoxyribonuclease, esterase, alpha-galactosidase, beta-galactosidase, glucoamylase, alpha-glucosidase, beta-glucosidase, invertase, laccase, lipase, mannosidase, mutanase, oxidase, a pectinolytic enzyme, peroxidase, phytase, polyphenoloxidase, proteolytic enzyme, ribonuclease, transglutaminase or xylanase. The nucleic acid sequence is contained in the chromosome of the *Bacillus* cell. The *Bacillus* host cell is *Bacillus alkalophilus*, *Bacillus amyloliquefaciens*, *Bacillus brevis*, *Bacillus brevis*, *Bacillus circulans*, *Bacillus clausii*, *Bacillus coagulans*, *Bacillus firmus*, *Bacillus lautus*, *Bacillus lentus*, *Bacillus licheniformis*, *Bacillus megaterium*, *Bacillus pumilus*, *Bacillus stercorophilus*, *Bacillus subtilis*, or *Bacillus thuringiensis*. This method alternatively comprises cultivating a *Bacillus* cell in a medium conducive for the production of the polypeptide, where the *Bacillus* cell comprises a nucleic acid construct comprising a consensus promoter having the sequence TTGACA for the -35 region and TATAAT for the -10 region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and an mRNA processing/stabilizing sequence located downstream of the consensus promoter and upstream of the nucleic acid sequence encoding the polypeptide; and isolating the polypeptide from the cultivation medium. The consensus promoter is obtained from any bacterial promoter, preferably a *Bacillus* promoter. Preferred Cell: The *Bacillus* cell comprises a nucleic acid construct that further comprises a selectable marker gene. The cell can also contain no selectable marker gene.

USE - The methods are useful for producing a polypeptide in a *Bacillus* cell, and for producing a selectable marker-free mutant of a *Bacillus* cell.

EXAMPLE - No relevant example given. (57 pages)

L14 ANSWER 4 OF 6 HCAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 2001:630833 HCAPLUS

DOCUMENT NUMBER: 135:209984

TITLE: Manufacture of large crystals of  $\delta$ -endotoxins with *Bacillus thuringiensis* by increasing gene copy number

INVENTOR(S): Adams, Lee Fremont; Thomas, Michael David; Sloma, Alan P.; Widner, William R.

PATENT ASSIGNEE(S): Valant Biosciences, Inc., USA; Libertyville, Inc.

SOURCE: U.S., 19 pp., Cont.-in-part of U. S. Ser. No. 92,338, abandoned.

CODEN: USXXAM

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 4

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
US 6280720	B1	20010828	US 1994-274608	19940713
IN 178296	A	19970322	IN 1994-MA623	19940712
CA 2167178	AA	19950126	CA 1994-2167178	19940714
ZA 9405138	A	19950223	ZA 1994-5138	19940714

CZ 290015	B6	20020515	CZ 1996-96	19940714
ES 2196030	T3	20031216	ES 1994-922566	19940714
US 6280721	B1	20010828	US 1995-377891	19950125
US 6303382	B1	20011016	US 1996-771190	19961220
US 6270760	B1	20010807	US 1997-872571	19970610
US 5955367	A	19990921	US 1998-60288	19980414
PRIORITY APPLN. INFO.:			US 1993-92338	B2 19930715
			DK 1989-6396	A 19891218
			US 1992-853701	B2 19920526
			US 1994-274608	A2 19940713
			US 1995-377892	B1 19950125
			US 1995-378236	B1 19950125

AB The invention relates to a method for producing an integrant(s) of *Bacillus thuringiensis* which produces a larger quantity of a crystal  $\delta$ -endotoxin with a greater pesticidal activity than the crystal  $\delta$ -endotoxin produced by the corresponding parental strain. The crystal  $\delta$ -endotoxin produced by the integrant *Bacillus thuringiensis* will have an activity directed towards the same pest(s) as its parent *Bacillus thuringiensis* crystal  $\delta$ -endotoxin. The invention further relates to such integrants, compns. comprising such integrants, as well as methods for controlling a pest(s) using these compns. Integration is achieved by transformation with a plasmid that does not carry a replicon that functions in *B. thuringiensis* but that does carry sequence that will direct efficient integration of the plasmid into the host chromosome. Use of the *cryIII* and *cryIIIA* genes is demonstrated. Parasporal crystals from integrant hosts were up to twice as long and 30% wider than crystals from parental strains. LC50's for the  $\delta$ -endotoxins of the integrants were 3-8-fold lower than those of the parental strains.

REFERENCE COUNT: 47 THERE ARE 47 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L14 ANSWER 5 OF 6 HCAPLUS COPYRIGHT 2006 ACS on STN

ACCESSION NUMBER: 1999:566201 HCAPLUS

DOCUMENT NUMBER: 131:180803

TITLE: Nucleic acid vectors for recombinant production of heterologous proteins in a *Bacillus* cell

INVENTOR(S): Widner, William; Sloma, Alan; Thomas, Michael D.

PATENT ASSIGNEE(S): Novo Nordisk Biotech, Inc., USA

SOURCE: PCT Int. Appl., 90 pp.

CODEN: PIXXD2

DOCUMENT TYPE: Patent

LANGUAGE: English

FAMILY ACC. NUM. COUNT: 3

PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
-----	----	-----	-----	-----
WO 9943835	A2	19990902	WO 1999-US4360	19990226
WO 9943835	A3	19991125		
W:	AL, AU, BB, BG, BR, CA, CN, CU, CZ, EE, GE, HU, IL, IN, IS, JP, KP, KR, LC, LK, LR, LT, LU, LV, MG, MK, MN, MX, NO, NZ, PL, RO, SG, SI, SK, TR, TT, UA, UZ, VN, YU, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			
RW:	GH, GM, KE, LS, MW, SD, SL, SZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG			
US 5955310	A	19990921	US 1998-31442	19980226
AU 9929756	A1	19990915	AU 1999-29756	19990226
EP 1056873	A2	20001206	EP 1999-911012	19990226
R:	AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, NL, SE, PT, IE, FI			
JP 2002504379	T2	20020212	JP 2000-533574	19990226
CN 1510145	A	20040707	CN 2003-2003158121	19990226
US 2003170876	A1	20030911	US 2001-834271	20010412

PRIORITY APPLN. INFO.:

US 1998-31442	A 19980226
US 1999-256377	B3 19990224
WO 1999-US4360	W 19990226

AB The present invention relates to methods for producing a polypeptide, comprising: (a) cultivating a *Bacillus* host cell in a medium conducive for the production of the polypeptide, wherein the *Bacillus* cell comprises a nucleic acid construct comprising (i) a tandem promoter in which each promoter sequence of the tandem promoter is operably linked to a single copy of a nucleic acid sequence encoding the polypeptide, and alternatively also (ii) an mRNA processing/stabilizing sequence located downstream of the tandem promoter and upstream of the nucleic acid sequence encoding the polypeptide; and (b) isolating the polypeptide from the cultivation medium. The present invention also relates to methods for producing a polypeptide, comprising: (a) cultivating a *Bacillus* host cell in a medium conducive for the production of the polypeptide, wherein the *Bacillus* cell comprises a nucleic acid construct comprising (i) a "consensus" promoter having the sequence TTGACA for the "-35" region and TATAAT for the "-10" region operably linked to a single copy of a nucleic acid sequence encoding the polypeptide and (ii) an mRNA processing/stabilizing sequence located downstream of the "consensus" promoter and upstream of the nucleic acid sequence encoding the polypeptide; and (b) isolating the polypeptide from the cultivation medium. Random promoters are created by placing promoters such as amyQ and amyL upstream of the cryIIIA promoter and its mRNA stabilizing sequence. Alternatively, "consensus" amyQ promoters are created with the cryIIIA mRNA stabilizing sequence, as well as tandem copies of a single promoter such as the short consensus amyQ dimer and trimer promoters. All of these approaches lead to significantly higher levels of SAVINASE gene expression (up to 620%) in *Bacillus* cells when compared to the levels obtained using single promoters such as amyQ and amyL.

L14 ANSWER 6 OF 6 BIOTECHDS COPYRIGHT 2006 THE THOMSON CORP. on STN  
DUPLICATE 2

ACCESSION NUMBER: 1995-04013 BIOTECHDS

TITLE: Methods for producing large *Bacillus thuringiensis* crystals; crystal protein overproduction by integration via homologous recombination and marker rescue, for use in biological control agent strain improvement

AUTHOR: Adams L F; Thomas M D; Sloma A P; Widner W  
R

PATENT ASSIGNEE: Novo-Nordisk-Entotech; Novo-Nordisk-Biotech

PATENT INFO: WO 9502695 26 Jan 1995

APPLICATION INFO: WO 1994-US7955 14 Jul 1994

PRIORITY INFO: US 1993-92338 15 Jul 1993

DOCUMENT TYPE: Patent

LANGUAGE: English

OTHER SOURCE: WPI: 1995-067339 [09]

AB A new integrant of *Bacillus thuringiensis* produces an increased quantity of delta-endotoxin crystal protein with improved pesticidal activity against the same pest as the parent, and is produced by homologous recombination using a vector with a homologous region and a selectable marker (e.g. antibiotic-resistance) in the presence of increasing amounts of a selective agent for gene amplification. The toxin is encoded by the cryI, cryII, cryIII, cryIV, cryV or cryVI gene, preferably cryIIIA. The integrant may be *B. thuringiensis* EMCC 0082 (NRRL B-21106), EMCC 0083 (NRRL B-21107), EMCC 0115 (NRRL B-21286) or EMCC 00116 (NRRL B-21287). The toxin gene is from *B. thuringiensis* subsp. *kurstaki*, *aizawai* (preferred), *galleriae*, *entomocidus*, *tenebrionis* (preferred), *thuringiensis*, *alesti*, *canadiensis*, *darmstadiensis*, *dendrolimus*, *finitimus*, *kenyae*, *morrisoni*, *subtoxicus*, *toumanoffi* or *israelensis*. A 2-vector double selectable marker rescue system may be used for recombination. The host is incubated at 37 deg. The crystal protein-producing strain is active against Coleoptera or Lepidoptera



insect pests. (44pp)

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(FILE 'HOME' ENTERED AT 14:23:15 ON 23 FEB 2006)

FILE 'MEDLINE, EMBASE, BIOSIS, BIOTECHDS, SCISEARCH, HCAPLUS, NTIS, LIFESCI' ENTERED AT 14:23:47 ON 23 FEB 2006

L1	395086 S BACILLUS
L2	560 S CRYIIIA
L3	516 S L1 AND L2
L4	635 S "TTGACA" OR "TATAAT"
L5	13 S L3 AND L4
L6	6 DUP REM L5 (7 DUPLICATES REMOVED)
L7	7 S L2 AND "CONSENSUS PROMOTER"
L8	5 DUP REM L7 (2 DUPLICATES REMOVED)
	E WIDNER W/AU
L9	119 S E3-E9
	E SLOMA A/AU
L10	124 S E3
	E THOMAS M D/AU
L11	404 S E3
L12	632 S L9 OR L10 OR L11
L13	8 S L2 AND L12
L14	6 DUP REM L13 (2 DUPLICATES REMOVED)

	Issue Date	Pages	Document ID	Title
1	20051208	21	US 20050272695 A1	Fast dissolving dried hyaluronic acid product
2	20051201	21	US 20050267068 A1	Dried and agglomerated hyaluronic acid product
3	20051006	94	US 20050221446 A1	Methods for producing hyaluronic acid in a Bacillus cell
4	20031002	22	US 20030186380 A1	Methods for producing secreted polypeptides having L-asparaginase activity
5	20030911	57	US 20030170876 A1	Methods for producing a polypeptide in a bacillus cell
6	20010703	54	US 6255076 B1	Methods for producing a polypeptide in a Bacillus cell

	Issue Date	Page s	Document ID	Title
1	20001031	35	US 6140104 A	Nucleotide sequences for the control of the expression of DNA sequences in a cell host

	Issue Date	Pages	Document ID	Title
1	20060223	32	US 2006004028 1 A1	Polyhydroxybutyrate polymerase
2	20051208	21	US 2005027269 5 A1	Fast dissolving dried hyaluronic acid product
3	20051201	21	US 2005026706 8 A1	Dried and agglomerated hyaluronic acid product
4	20051103	114	US 2005024679 8 A1	Novel glyphosate-N-acetyltransferase (GAT) genes
5	20051006	94	US 2005022144 6 A1	Methods for producing hyaluronic acid in a Bacillus cell
6	20050901	47	US 2005019162 8 A1	Antibiotics based upon bacteriophage lysis proteins
7	20050414	55	US 2005007961 7 A1	Glucose transport mutants for production of biomaterial
8	20050324	34	US 2005006456 5 A1	POLYHYDROXYBUTYRATE POLYMERASE
9	20040819	16	US 2004016241 7 A1	Protease, a gene therefor and the use thereof
10	20040429	86	US 2004008277 0 A1	Novel glyphosate N-acetyltransferase (GAT) genes
11	20040226	108	US 2004003826 2 A1	Ribulose 1,5-bisphosphate carboxylase/oxygenase polypeptides and related polynucleotides
12	20040205	218	US 2004002320 5 A1	Method of recovering a nucleic acid encoding a proteinaceous binding domain which binds a target material

	Issue Date	Pages	Document ID	Title
13	20040108	218	US 2004000553 9 A1	Nucleic acids, genetic constructs, and library of nucleic acids encoding fusion proteins
14	20031218	61	US 2003023240 6 A1	Bacterial strains which overproduce riboflavin
15	20031204	114	US 2003022524 9 A1	32-kDa protein derived from mycobacterium tuberculosis and related peptides
16	20031127	197	US 2003021988 6 A1	Directed evolution of novel binding proteins
17	20031127	224	US 2003021972 2 A1	Fusion proteins, modified filamentous bacteriophage, and populations or libraries of same
18	20031002	22	US 2003018638 0 A1	Methods for producing secreted polypeptides having L-asparaginase activity
19	20030918	142	US 2003017590 2 A1	Methods for producing hyaluronan in a recombinant host cell
20	20030911	57	US 2003017087 6 A1	Methods for producing a polypeptide in a bacillus cell
21	20030619	43	US 2003011562 7 A1	Coniothyrium minitans beta-(1,3) exoglucanase gene cbeg1
22	20030619	190	US 2003011371 7 A1	Directed evolution of novel binding proteins
23	20030501	139	US 2003008348 0 A1	Novel glyphosate N-acetyl transferase (GAT) genes

24	20021017	203	US 2002015088 1 A1	Directed evolution of novel binding proteins
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	Issue Date	Pages	Document ID	Title
25	20020912	33	US 20020127240 A1	Leptospiral rare outer membrane proteins
26	20060103	70	US 6982314 B2	Lawsonia intracellularis proteins, and related methods and materials
27	20051227	303	US 6979538 B2	Directed evolution of novel binding proteins
28	20050419	36	US 6881560 B2	Polyhydroxybutyrate polymerase
29	20050111	56	US 6841376 B2	Bistable genetic toggle switch
30	20040511	42	US 6734344 B2	Coniothyrium minitans .beta.-(1,3) exoglucanase gene cbeg 1
31	20040413	217	US 6720488 B2	Transgenic maize seed and method for controlling insect pests
32	20040302	33	US 6699482 B2	Leptospira rare outer membrane proteins
33	20030422	60	US 6551813 B1	Nutrient medium for bacterial strains which overproduce riboflavin
34	20030311	117	US 6531138 B1	Recombinant polypeptides and peptides, nucleic acids coding for the same and use of these polypeptides and peptides in the diagnostic of tuberculosis
35	20030304	35	US 6528706 B1	Polyhydroxybutyrate polymerase
36	20020521	32	US 6392126 B1	Adenosine deaminase homologues and uses thereof

37	20011127	109	US 6322995 B1	Riboflavin production
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	Issue Date	Pages	Document ID	Title
38	20011106	36	US 6313376 B1	Maize aquaporins and uses thereof
39	20010918	11	US 6291230 B1	Galk promoter
40	20010724	36	US 6265636 B1	Pyruvate dehydrogenase kinase polynucleotides, polypeptides and uses thereof
41	20010703	54	US 6255076 B1	Methods for producing a polypeptide in a Bacillus cell
42	20010626	21	US 6252140 B1	Promoters from chlorella virus genes providing for expression of genes in prokaryotic and eukaryotic hosts
43	19990720	63	US 5925538 A	Bacterial strains which overproduce riboflavin
44	19990629	119	US 5916558 A	Recombinant polypeptides and peptides, nucleic acids coding for the same and use of these polypeptides and peptides in the diagnostic of tuberculosis
45	19981201	69	US 5843426 A	Salmonella vaccines
46	19981117	62	US 5837528 A	Bacterial strains which overproduce riboflavin
47	19981117	181	US 5837500 A	Directed evolution of novel binding proteins
48	19981117	33	US 5837263 A	Leptospira membrane proteins
49	19980825	25	US 5798235 A	Gene encoding bacterial acetoacetylco a reductase

50	19980811	21	US 5792924 A	Biologically safe plant transformation system
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	Issue Date	Page s	Document ID	Title
51	19980324	14	US 5731151 A	Regulator of contact-mediated hemolysin
52	19970902	32	US 5663063 A	Method for producing polyester biopolymers
53	19970826	24	US 5661026 A	Gene encoding bacterial beta-ketothiolase
54	19961105	176	US 5571698 A	Directed evolution of novel binding proteins
55	19960709	33	US 5534432 A	Polyhydroxybutyrate polymerase
56	19960430	25	US 5512669 A	Gene encoding bacterial acetoacetyl-CoA reductase
57	19960116	34	US 5484718 A	Nodulation gene promoter
58	19960109	21	US 5482852 A	Biologically safe plant transformation system
59	19950404	199	US 5403484 A	Viruses expressing chimeric binding proteins
60	19931005	33	US 5250430 A	Polyhydroxyalkanoate polymerase
61	19930914	32	US 5245023 A	Method for producing novel polyester biopolymers
62	19930720	25	US 5229279 A	Method for producing novel polyester biopolymers
63	19930706	19	US 5225341 A	Biologically safe plant transformation system using a Ds transposon
64	19921215	20	US 5171673 A	Expression of heterologous DNA using the bacillus coagulans amylase gene

65	19910122	13	US 4987078 A	Plasmid vectors for expression in Escherichia coli and/or Bacillus subtilis
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	Issue Date	Page s	Document ID	Title
66	19900417	36	US 4918006 A	Gene coding for insecticidal crystal protein
67	19880906	42	US 4769327 A	Secretion vector

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<b>1</b>	L1	1	"6852518".pn.
<b>2</b>	L2	4	"GST-40"
<b>3</b>	L3	1	l1 and l2
<b>4</b>	L4	4140 1	bacillus
<b>5</b>	L5	365	cryIIIA
<b>6</b>	L6	314	l4 same l5
<b>7</b>	L7	495	"TATAAT" or "TTGACA"
<b>8</b>	L8	6	l6 same l7
<b>9</b>	L9	3016	"downstream region"
<b>10</b>	L10	1	l6 same l9
<b>11</b>	L11	4113 91	WIDNER SLOMA THOMAS
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